

U.S. Appln. No. 08/887,505  
Our Ref. No.: HYZ-040CIP/ 47508.250US3  
Reply To Office Action of October 20, 2005

**APPENDIX A**

Attached is a marked-up copy of pages 29-30 of the application pursuant to 37 C.F.R. § 1.121(b).

TABLE 1E

Contiguous and Non-contiguous Multiplex Oligonucleotides					
SEQ ID NO:	Oligo	Sequence	Duplex target	Triplex Target (Purine Strand)	Description
159	HCV-222	CCCUCCCCGGGG-Y-tcctg	-218 to -227	-212 to -222	
160	HCV-223	GGGGG-Y-tcctg	-218 to -227	None	
161	HCV-224	CCUCCCCC-Y-(GGGGG)-tcctg	-218 to -227	-212 to -222	( )= branched triplex forming sequence, 3'-5'
162	HCV-225	GGGGG-Y-tcctg	-218 to -227	None	
163	HCV-226	CCCUCCCCGGG-Y-(CCCCC)-tcctg	-218 to -227	-212 to -222	
164	HCV-227	GGGGG-Y-(CCCCC)-tcctg	-218 to -227	-212 to -222	( )= branched triplex forming sequence, 3'-5'
165	HCV-228	CCCUCCCCGGGG-Y-tcctg	-218 to -227	-212 to -217	
166	HCV-229	GUCUACGAGAGGGGG-Y-(CCCCCCCUCCC)-tcctg	-218 to -227/ -18 to -9	-212 to -222	( )= branched triplex forming sequence, 3'-5'



167	HCV-230	GUUCUACGAGGGGG-Y-tcctg	-218 to -227/ -18 to -9	None	
168	HCV-231	GUUCUACGAGGGGG-tcctg	-218 to -227/ -18 to -9	None	
169	HCV-232	GUUCUACGAGA-Y-(CCUCCG)-gggggg	-218 to -222/ -18 to -9	-212 to -217	( )=branched triples triplex- forming sequence, 3'-5'
170	HCV-233	GUUCUACGAGA-Y-ggggg	-218 to -222/ -18 to -9	None	
171	HCV-234	CCCGGGAGGGGGGG-Y- (CCCCCCCCUCCC)-tcctg	-209 to -227	-212 to -222	( )=branched triples triplex- forming sequence, 3'-5'
172	HCV-235	CCCGGGAGGGGGGG-Y-tcctg	-209 to -227	None	

Upper case = 2'-OMe RNA

Lower case = DNA

Y = asymmetric branching monomer

Best Available Copy

